



Exhibit C

Blast SEQ ID NO:1 versus human genomic

MEGABLAST 1.2.3-Paracel [2001-11-20]

Reference:

Zheng Zhang, Scott Schwartz, Lukas Wagner, and Webb Miller (2000),
"A greedy algorithm for aligning DNA sequences",
J Comput Biol 2000; 7(1-2):203-14.
Database: Homo_sapiens.latestgp.masked.fa
44,521 sequences; 200,768,834,160 total letters

Query= LEX161seqid1
(2115 letters)

Sequences producing significant alignments:	Score	E
Value	(bits)	
AC016643.6.1.179969	734	0.0
AC008958.7.1.109526	520	e-
144		

>AC016643.6.1.179969
Length = 179969

Score = 734 bits (370), Expect = 0.0
Identities = 370/370 (100%)
Strand = Plus / Plus

Query: 957 agatgatttaagccagaaggttaaaacccttgggtgaagcagaacgagagtttattttgaa
1016

|||||
Sbjct: 60550 agatgatttaagccagaaggttaaaacccttgggtgaagcagaacgagagtttattttgaa
60609

Query: 1017 tttgaagaaaaaggaatgcaaagacaggggttttgaatatgatgggaaaatcaatgcctg
1076

|||||
Sbjct: 60610 tttgaagaaaaaggaatgcaaagacaggggttttgaatatgatgggaaaatcaatgcctg
60669

Query: 1077 ggatctatattactacatgactcagacagaggaactcaagtattccatagaccaagagtt
1136

|||||
Sbjct: 60670 ggatctatattactacatgactcagacagaggaactcaagtattccatagaccaagagtt
60729

Query: 1137 cctcaaggaatacttcccaattgaggtgggtcactgaaggcttgctgaacacctaccagga
1196

|||||
Sbjct: 60730 cctcaaggaatacttcccaattgaggtgggtcactgaaggcttgctgaacacctaccagga
60789

Query: 1197 gttggtgggactttcatttgaacaaatgacagatgctcatgtttggaacaagagtgttac
1256
|||||
Sbjct: 60790 gttggtgggactttcatttgaacaaatgacagatgctcatgtttggaacaagagtgttac
60849

Query: 1257 actttatactgtgaaggataaagctacaggagaagtattgggacagttctatttggacct
1316
|||||
Sbjct: 60850 actttatactgtgaaggataaagctacaggagaagtattgggacagttctatttggacct
60909

Query: 1317 ctatccaagg 1326
|||||
Sbjct: 60910 ctatccaagg 60919

Score = 520 bits (262), Expect = e-144
Identities = 262/262 (100%)
Strand = Plus / Plus

Query: 40 agagttggtggtccaggattttactcagaatgacgtaggaagagaagtgatgtctcct
99
|||||
Sbjct: 30994 agagttggtggtccaggattttactcagaatgacgtaggaagagaagtgatgtctcct
31053

Query: 100 cttcaggcaatgtcttcctatactgtggctggcagaaatgttttaagatgggatctttca
159
|||||
Sbjct: 31054 cttcaggcaatgtcttcctatactgtggctggcagaaatgttttaagatgggatctttca
31113

Query: 160 ccagagcaaattaaaacaagaactgaggagctcattgtgcagaccaaacaggtgtacgat
219
|||||
Sbjct: 31114 ccagagcaaattaaaacaagaactgaggagctcattgtgcagaccaaacaggtgtacgat
31173

Query: 220 gctgttggaatgctcgggtattgaggaagtaacttacgagaactgtctgcaggcactggca
279
|||||
Sbjct: 31174 gctgttggaatgctcgggtattgaggaagtaacttacgagaactgtctgcaggcactggca
31233

Query: 280 gatgtagaagtaaagtatatag 301
|||||
Sbjct: 31234 gatgtagaagtaaagtatatag 31255

Score = 405 bits (204), Expect = e-110

Identities = 204/204 (100%)
Strand = Plus / Plus

Query: 1324 agggaaggaaaataacaatcatgcggcctgcttcggtctccagcctggctgccttctgcct
1383

|||||
Sbjct: 64886 agggaaggaaaataacaatcatgcggcctgcttcggtctccagcctggctgccttctgcct
64945

Query: 1384 gatggaagccggatgatggcagtggtgccctcgtggtgaacttctcacagccagtggca
1443

|||||
Sbjct: 64946 gatggaagccggatgatggcagtggtgccctcgtggtgaacttctcacagccagtggca
65005

Query: 1444 ggtcgtccctctctcctgagacacgacgaggtgaggacttactttcatgagtttggtcac
1503

|||||
Sbjct: 65006 ggtcgtccctctctcctgagacacgacgaggtgaggacttactttcatgagtttggtcac
65065

Query: 1504 gtgatgcatcagatttgtgcacag 1527

|||||
Sbjct: 65066 gtgatgcatcagatttgtgcacag 65089

Score = 379 bits (191), Expect = e-102
Identities = 191/191 (100%)
Strand = Plus / Plus

Query: 1526 agactgattttgcacgatttagcggaacaaatgtggaaactgactttgtagaggtgccat
1585

|||||
Sbjct: 81947 agactgattttgcacgatttagcggaacaaatgtggaaactgactttgtagaggtgccat
82006

Query: 1586 cgcaaatgcttgaaaattgggtgtgggacgtcgattccctccgaagattgtcaaaacatt
1645

|||||
Sbjct: 82007 cgcaaatgcttgaaaattgggtgtgggacgtcgattccctccgaagattgtcaaaacatt
82066

Query: 1646 ataaagatggaagccctattgcagacgatctgcttgaaaaacttggtgcttctaggctgg
1705

|||||
Sbjct: 82067 ataaagatggaagccctattgcagacgatctgcttgaaaaacttggtgcttctaggctgg
82126

Query: 1706 tcaacacaggt 1716

|||||
Sbjct: 82127 tcaacacaggt 82137

Score = 323 bits (163), Expect = 2e-85
Identities = 163/163 (100%)
Strand = Plus / Plus

Query: 661 ggtgctcttcctgatgatttcattgacagtttagaaaagacagatgatgacaagtataaa
720
|||||
Sbjct: 53694 ggtgctcttcctgatgatttcattgacagtttagaaaagacagatgatgacaagtataaa
53753

Query: 721 attaccttaaaatatccacactatttcctgtcatgaagaaatggtgtatccctgaaacc
780
|||||
Sbjct: 53754 attaccttaaaatatccacactatttcctgtcatgaagaaatggtgtatccctgaaacc
53813

Query: 781 agaagaaggatggaaatggcttttaatacaagggtgcaaagagg 823
|||||
Sbjct: 53814 agaagaaggatggaaatggcttttaatacaagggtgcaaagagg 53856

Score = 301 bits (152), Expect = 8e-79
Identities = 152/152 (100%)
Strand = Plus / Plus

Query: 300 agtggaaaggaccatgctagactttccccagcatgtatcctctgacaaagaagtacgagc
359
|||||
Sbjct: 35386 agtggaaaggaccatgctagactttccccagcatgtatcctctgacaaagaagtacgagc
35445

Query: 360 agcaagtacagaagcagacaaaagactttctcgttttgatattgagatgagcatgagagg
419
|||||
Sbjct: 35446 agcaagtacagaagcagacaaaagactttctcgttttgatattgagatgagcatgagagg
35505

Query: 420 agatatatttgagagaattgttcatttacagg 451
|||||
Sbjct: 35506 agatatatttgagagaattgttcatttacagg 35537

Score = 280 bits (141), Expect = 3e-72
Identities = 141/141 (100%)
Strand = Plus / Plus

Query: 1842 aggcacaaatatgccagctacctttggacatttggcagggggatacgatggccaatatta
1901
|||||

Sbjct: 84687 aggcacaaatatgccagctacctttggacatttggcagggggatacgatggccaatatta
84746

Query: 1902 tggatatctttggagtgaagtattttccatggatatgttttacagctgttttaaaaaaga
1961

|||||
Sbjct: 84747 tggatatctttggagtgaagtattttccatggatatgttttacagctgttttaaaaaaga
84806

Query: 1962 agggataatgaatccagaggt 1982

|||||
Sbjct: 84807 agggataatgaatccagaggt 84827

Score = 270 bits (136), Expect = 3e-69
Identities = 137/138 (99%)
Strand = Plus / Plus

Query: 821 aggaaaacaccataattttgcagcagctactcccactgcgaaccaaggtggccaaactac
880

|||||
Sbjct: 58207 aggaaaacaccataattttgcagcagctactcccactgcgaaccaaggtggccaaactac
58266

Query: 881 tcggttatagcacacatgctgacttcgctccttgaaatgaacactgcaaagagcacaagcc
940

|||||
Sbjct: 58267 tcggttatagcacacatgctgacttcgctccttgaaatgaacactgcaaagagcacaagcc
58326

Query: 941 gcgtaacagcytttctag 958

|||||
Sbjct: 58327 gcgtaacagcctttctag 58344

Score = 268 bits (135), Expect = 1e-68
Identities = 136/137 (99%)
Strand = Plus / Plus

Query: 1979 aggttggaatgaaatacagaaacctaatacctgaaacctgggggatctctggacggcatgg
2038

|||||
Sbjct: 95214 aggttggaatgaaatacagaaacctaatacctgaaacctgggggatctctggacggcatgg
95273

Query: 2039 acatgctccacaatttcttgaaacgtgagccaaacaaaaagcgttcctaatagagtagag
2098

|||||
Sbjct: 95274 acatgctccacaatttcttgaaacgtgagccaaacaaaaagcgttcctaatagagtagag
95333

Query: 2099 gcctgcatgctycgtga 2115
|||||||
Sbjct: 95334 gcctgcatgctccgtga 95350

Score = 262 bits (132), Expect = 7e-67
Identities = 132/132 (100%)
Strand = Plus / Plus

Query: 1713 aggtcttctgaccctgcgccagattgttttgagcaaagttgatcagtctcttcatac
1772
|||||||
Sbjct: 82469 aggtcttctgaccctgcgccagattgttttgagcaaagttgatcagtctcttcatac
82528

Query: 1773 cacatcgctggatgctgcaagtgaatatgccaaatactgctcagaaatattaggagttgc
1832
|||||||
Sbjct: 82529 cacatcgctggatgctgcaagtgaatatgccaaatactgctcagaaatattaggagttgc
82588

Query: 1833 agctactccagg 1844
|||||||
Sbjct: 82589 agctactccagg 82600

Score = 222 bits (112), Expect = 6e-55
Identities = 116/117 (99%), Gaps = 1/117 (0%)
Strand = Plus / Plus

Query: 443 atttacaggaaacctgtgatctggggaagataaaacctgaggccagacgataacttgaaa
502
|||||||
Sbjct: 49854 attt-caggaaacctgtgatctggggaagataaaacctgaggccagacgataacttgaaa
49912

Query: 503 agtcaattaaaatggggaaaagaaatgggctccatcttcctgaacaagtacagaatg 559
|||||||
Sbjct: 49913 agtcaattaaaatggggaaaagaaatgggctccatcttcctgaacaagtacagaatg
49969

Score = 208 bits (105), Expect = 1e-50
Identities = 105/105 (100%)
Strand = Plus / Plus

Query: 559 gaaatcaaataatgaagaaaagaatgagtgagctatgtattgattttaacaaaaacctc
618
|||||||
Sbjct: 52613 gaaatcaaataatgaagaaaagaatgagtgagctatgtattgattttaacaaaaacctc
52672

Query: 619 aatgaggatgataccttccttgatattttccaaggctgaacttggt 663
|||||
Sbjct: 52673 aatgaggatgataccttccttgatattttccaaggctgaacttggt 52717

>AC008958.7.1.109526
Length = 109526

Score = 520 bits (262), Expect = e-144
Identities = 262/262 (100%)
Strand = Plus / Minus

Query: 40 agagttggtggttccaggattttactcagaatgacgtaggaagagaagtgatgtctcct
99
|||||
Sbjct: 24199 agagttggtggttccaggattttactcagaatgacgtaggaagagaagtgatgtctcct
24140

Query: 100 cttcaggcaatgtcttcctatactgtggctggcagaaatgttttaagatgggatctttca
159
|||||
Sbjct: 24139 cttcaggcaatgtcttcctatactgtggctggcagaaatgttttaagatgggatctttca
24080

Query: 160 ccagagcaaattaaaacaagaactgaggagctcattgtgcagaccaaacaggtgtacgat
219
|||||
Sbjct: 24079 ccagagcaaattaaaacaagaactgaggagctcattgtgcagaccaaacaggtgtacgat
24020

Query: 220 gctgttggaatgctcggatttgaggaagtaacttacgagaactgtctgcaggcactggca
279
|||||
Sbjct: 24019 gctgttggaatgctcggatttgaggaagtaacttacgagaactgtctgcaggcactggca
23960

Query: 280 gatgtagaagtaaagtatatag 301
|||||
Sbjct: 23959 gatgtagaagtaaagtatatag 23938

Score = 323 bits (163), Expect = 2e-85
Identities = 163/163 (100%)
Strand = Plus / Minus

Query: 661 ggtgctcttcctgatgatttcattgacagtttagaaaagacagatgatgacaagtataaa
720
|||||
Sbjct: 1504 ggtgctcttcctgatgatttcattgacagtttagaaaagacagatgatgacaagtataaa
1445

Query: 721 attaccttaaaatatccacactatttcctgtcatgaagaaatgttgatccctgaaacc
780

|||||
Sbjct: 1444 attaccttaaaatatccacactatttcctgtcatgaagaaatgttgatccctgaaacc
1385

Query: 781 agaagaaggatggaaatggcttttaatacaaggtgcaaagagg 823

|||||
Sbjct: 1384 agaagaaggatggaaatggcttttaatacaaggtgcaaagagg 1342

Score = 301 bits (152), Expect = 8e-79
Identities = 152/152 (100%)
Strand = Plus / Minus

Query: 300 agtggaaaggaccatgctagactttccccagcatgtatcctctgacaaagaagtacgagc
359

|||||
Sbjct: 19806 agtggaaaggaccatgctagactttccccagcatgtatcctctgacaaagaagtacgagc
19747

Query: 360 agcaagtacagaagcagacaaaagactttctcgttttgatattgagatgagcatgagagg
419

|||||
Sbjct: 19746 agcaagtacagaagcagacaaaagactttctcgttttgatattgagatgagcatgagagg
19687

Query: 420 agatatatttgagagaattgttcatttacagg 451

|||||
Sbjct: 19686 agatatatttgagagaattgttcatttacagg 19655

Score = 222 bits (112), Expect = 6e-55
Identities = 116/117 (99%), Gaps = 1/117 (0%)
Strand = Plus / Minus

Query: 443 atttacaggaaacctgtgatctggggaagataaaacctgaggccagacgatacttgaaa
502

|||||
Sbjct: 5344 attt-caggaaacctgtgatctggggaagataaaacctgaggccagacgatacttgaaa
5286

Query: 503 agtcaattaaaaatggggaaaagaaatgggctccatcttcctgaacaagtacagaatg 559

|||||
Sbjct: 5285 agtcaattaaaaatggggaaaagaaatgggctccatcttcctgaacaagtacagaatg 5229

Score = 208 bits (105), Expect = 1e-50
Identities = 105/105 (100%)
Strand = Plus / Minus

Query: 559 gaaatcaaataaatgaagaaaagaatgagtgagctatgtattgattttaacaaaaacctc
618

Sbjct: 2585 gaaatcaaataaatgaagaaaagaatgagtgagctatgtattgattttaacaaaaacctc
2526

Query: 619 aatgaggatgataccttccttgatattttccaaggctgaacttggt 663

Sbjct: 2525 aatgaggatgataccttccttgatattttccaaggctgaacttggt 2481

Score = 81.8 bits (41), Expect = 2e-12

Identities = 41/41 (100%)

Strand = Plus / Minus

Query: 1 atgatcgcccgggtgccttttggtgtgcaagcctccgcag 41

Sbjct: 60390 atgatcgcccgggtgccttttggtgtgcaagcctccgcag 60350